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Prevalence of Multi-Gastrointestinal Infections With Helminth, Protozoan and Campylobacter spp. in Guatemalan Children

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Prevalence of multi-gastrointestinal infections with helminth, protozoan and *Campylobacter* spp. in Guatemalan children

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Abstract

Background: The prevalence of multi-infections with helminthes, protozoans and *Campylobacter* spp. in Guatemalan children is a reflection of differences in the risk factors related to pathogen transmission.

Methodology: Two hundred and eighty-nine fecal samples were collected from children of the Guatemalan highlands and patterns of pathogen occurrences were evaluated using an immunoassay for *Campylobacter* spp., a formalin-ether concentration followed by observation of unstained slides for helminthes and trichome stains of fecal smears for protozoans. Specimens were examined microscopically using 100, 400 and 1000x magnification.

Results: Prevalence of *Ascaris lumbricoides*, *Campylobacter* spp., *Giardia duodenalis*, *Entamoeba histolytica*/E. *dispar* and *Trichuris trichiura* were 55.1%, 30.8%, 21.5%, 19.8% and 19.4%, respectively. Overall, the prevalence of at least one intestinal pathogen was 85.5%. Multi-infections were found in 43% of the children harboring pathogens.

Conclusions: Infections with *Campylobacter* spp., *E. histolytica*/E. *dispar*, *T. trichiura* and *G. duodenalis* were closely associated with the presence of co-infection with *A. lumbricoides*. *T. trichiura* infection was related to co-infection with *A. lumbricoides* and *Campylobacter* spp. Infections with *G. duodenalis* and *T. trichiura* were related to co-infections with either *Campylobacter* spp. or *E. histolytica*/E. *dispar*. The prevalence of multi-gastrointestinal infections with helminthes, protozoans and *Campylobacter* spp. in children was found to be related to age and gender.

Key words: Guatemala, diarrhoea, helminthes, protozoans, *Campylobacter*

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Introduction

Contaminated drinking water and soil and the lack of feces treatment are significant risk factors in the transmission of gastrointestinal pathogens [1,2]. Pathogens observed in multi-infections share the similar fecal-oral route of transmission, have similar age profiles, and are related to behavioral and occupational traits that increase their exposure to humans [3]. Under these conditions, multi-infections of helminths, protozoa and/or bacteria may be coincidental or enhance each other's ability to become established as pathogens in the gastrointestinal tract [4,5].

To better understand the risk factors associated with multi-gastrointestinal infections in children, a prevalence study was conducted in two communities in the Guatemalan highlands and three riparian communities. The study evaluated the prevalence of

multi-infections with *Ascaris lumbricoides*, *Trichuris trichiura*, *Giardia duodenalis*, *Entamoeba histolytica*/E. *dispar* and *Campylobacter* spp. in Guatemalan children.

Materials and methods

Characteristics of the communities studied

The study area included two communities in Guatemala, *i.e.*, Santa Maria de Jesus and La Mano de Leon, which are located at the 4,500 m level in the mountains around Antigua and three riparian communities, El Esol, Chichipute, and Rio Dulce, situated on the banks of the Rio Dulce River. The five communities were chosen at random from the surrounding villages. Fecal samples were obtained during patient visits to field clinics. Each village was suspect for soil-transmitted gastrointestinal helminthes and water-borne protozoans and bacteria.

Table 1. Prevalence of single pathogens in children.

	Number infected (%) Age Groups (yrs)				
	0-4	5-8	9-12	Total	
Male Pathogens					
<i>A. lumbricoides</i>	12 (8.57) ^a	6 (4.29)	5 (3.57)	23 (16.43)	
<i>T. trichuris</i>	1 (0.71)	4 (2.86)	0 (0.00)	5 (3.57)	
<i>E. histolytica/E. dispar</i>	2 (1.43)	1 (0.71)	0 (0.00)	3 (2.14)	
<i>G. lamblia</i>	9 (6.43)	5 (3.57)	0 (0.00)	14 (10.00)	
<i>Campylobacter</i> spp.	13 (9.29)	2 (1.43)	1 (0.71)	16 (11.43)	
Total	37 (26.43) ^b	18 (12.86) ^b	6 (4.29)	61 (43.57)	
Female Pathogens					All
<i>A. lumbricoides</i>	17 (12.14) ^a	11 (7.86) ^{a, c}	12 (8.57) ^{a, c}	40 (28.57) ^c	63 (45.00)
<i>T. trichuris</i>	1 (0.71)	1 (0.71)	0 (0.00)	2 (1.43)	7 (5.00)
<i>E. histolytica/E. dispar</i>	5 (3.57)	4 (2.86)	1 (0.71)	10 (7.14) ^c	13 (9.29)
<i>G. lamblia</i>	6 (4.29)	4 (2.86)	0 (0.00)	10 (7.14)	24 (17.14)
<i>Campylobacter</i> spp.	6 (4.29)	8 (5.71)	3 (2.14)	17 (12.14)	33 (23.57)
Total	36 (25.71) ^b	28 (20.00) ^{b, c}	16 (11.43) ^c	79 (56.43)	140 (100.00)

^a*A. lumbricoides* was significantly higher than other infections, $P < 0.05$.^bInfections were significantly higher in 0-4 and 5-8 yr-old males, $P < 0.05$.^cInfections in females were significantly higher than those in males.

In comparisons between data collected within each of the two highland and the three riparian communities, differences of multi-infection patterns could not be measured due to the small sample sizes of each age group. The two geographical areas were similar in that their economies were based on agricultural activities and they both had open sewer systems with poor sanitation practices and untreated drinking water. They differed with respect to diet: the riparian communities supplemented their diets with fish, which contrasted the mainly vegetable diet of the highland communities.

Diarrhoea specimens from 289 children aged 0 to 12 years old were collected and analyzed for pathogens and the results were pooled. Diarrhoeal stools were obtained from patients with abdominal symptoms in order to increase the chances of identifying *Campylobacter* spp. infections along with helminth and protozoan infections. Diarrhoea was defined as three or more loose stools in a 24-hour period. A limitation in the data collected was the fact that the 0-4 age group did not contain enough participants to be broken down into multiple groups of 4-6 months of age and thus represents the most heterogeneous group of patients in the study.

Examination of feces for pathogens

Fecal analysis was restricted to five pathogens (*A. lumbricoides*, *Campylobacter* spp., *E. histolytica/E. dispar*, *G. duodenalis* and *T. trichiura*) based on the availability of diagnostic tests. To determine the presence of protozoan parasites, a smear of each fecal sample was trichrome-stained and examined microscopically using 1000x magnification. Fecal preparations (formalin-ether concentrations) were prepared and examined for the presence of helminth eggs using 100x and 400x magnification. *Campylobacter* spp. antigens were identified using an enzyme-linked immunosorbent assay (ELISA; Remel, Lenexa, KS). No attempt was made to concurrently assess the prevalence of other bacteria and parasites.

Committee approval

The present study is part of a research project entitled "Survey of human pathogens in Guatemala" and was approved by the Institutional Review Board and the Health Professions Division Research Committee of Nova Southeastern University. Informed consent was obtained from each patient's parent or guardian and, following diagnosis, the children were treated for observed infections.

Table 2. Prevalence of two pathogens in children.

	Number infected (%) Age groups (yrs)				
Male co-pathogens	0-4	5-8	9-12	Total	all
<i>A.l.</i> + <i>E.h.</i>	1(1.35)	3(4.06)	1(1.35)	5(6.76)	
<i>A.l.</i> + <i>G.d.</i>	1(1.35)	1(1.35)	1(1.35)	3(4.06)	
<i>A.l.</i> + <i>C.spp.</i>	5(6.76)	2(2.70)	2(2.70)	9(12.16)	
<i>A.l.</i> + <i>T.t.</i>	2(2.70)	3(4.06)	2(2.70)	7(9.46)	
<i>T.t.</i> + <i>E.h.</i>	0(0.00)	1(1.35)	0(0.00)	1(1.35)	
<i>T.t.</i> + <i>G.d.</i>	1(1.35)	1(1.35)	0(0.00)	2(2.70)	
<i>T.t.</i> + <i>C.spp.</i>	0(0.00)	0(0.00)	0(0.00)	0(0.00)	
<i>G.d.</i> + <i>E.h.</i>	2(2.70)	1(1.35)	0(0.00)	3(4.06)	
<i>G.d.</i> + <i>C.spp.</i>	3(4.06)	3(4.06)	0(0.00)	6(8.11)	
<i>E.h.</i> + <i>C.spp.</i>	1(1.35)	0(0.00)	0(0.00)	1(1.35)	
Total	16 (21.62) ^a	15 (20.28) ^a	6 (8.10)	37 (50.0)	
Female co-pathogens	0-4	5-8	9-12	Total	all
<i>A.l.</i> + <i>E.h.</i>	1 (1.35)	4 (5.41)	6 (8.11)	11 (14.86)	16 (21.62) ^b
<i>A.l.</i> + <i>G.d.</i>	3 (4.06)	0 (0.00)	0 (0.00)	3 (4.06)	6 (8.11)
<i>A.l.</i> + <i>C.spp.</i>	6 (8.11)	2 (2.70)	1 (1.35)	9 (12.16)	18 (24.32) ^b
<i>A.l.</i> + <i>T.t.</i>	1 (1.35)	3 (4.06)	1 (1.35)	5 (6.76)	12 (16.22) ^b
<i>T.t.</i> + <i>E.h.</i>	0 (0.00)	0 (0.00)	0 (0.00)	0 (0.00)	1 (1.35)
<i>T.t.</i> + <i>G.d.</i>	0 (0.00)	0 (0.00)	0 (0.00)	0 (0.00)	2 (2.70)
<i>T.t.</i> + <i>C.spp.</i>	0 (0.00)	0 (0.00)	1 (1.35)	1 (1.35)	1 (1.35)
<i>G.d.</i> + <i>E.h.</i>	1 (1.35)	0 (0.00)	0 (0.00)	1 (1.35)	4 (5.41)
<i>G.d.</i> + <i>C.spp.</i>	2 (2.70)	0 (0.00)	0 (0.00)	2 (2.70)	8 (10.81)
<i>E.h.</i> + <i>C.spp.</i>	0 (0.00)	2 (2.70)	3 (4.06)	5 (6.76)	6 (8.11)
Total	14 (18.92)	11 (14.86)	12 (16.22)	37 (50.00)	74 (100.00)

(*A.l.*: *Ascaris lumbricoides*, *T.t.*: *Trichuris trichiura*, *E.h.*: *Entamoeba histolytica*/*E. dispar*, *G.d.*: *Giardia duodenalis*, *C.spp.*: *Campylobacter* species)

^aCo-infections were significantly higher in 0-4 and 5-8 yr-old males, $P < 0.05$.

^b*A. lumbricoides* + *E. histolytica*/*E. dispar*, *A. lumbricoides* + *Campylobacter* sp. and *A. lumbricoides* + *T. trichiura* were the most prevalent co-infections in males and females, $P < 0.05$.

Statistical analysis

Adjusted odds ratios and 95% confidence intervals were computed for estimates of associations between enteropathogens, and McNemar's test was used to compute P-values between co-infections of the samples [6].

Results

Pathogens were identified from 247 (85.5%) of 289 diarrhoeal stools. One hundred and thirty-six specimens contained *A. lumbricoides* (55.1%), 76 *Campylobacter* spp. (30.8%), 53 *G. duodenalis* (21.5%), 49 *E. histolytica*/*E. dispar* (19.8%) and 48 *T. trichiura* (19.4%). Of the 247 infected patients, 140 had only one pathogen; 70 with helminthes (50.0%), 37 with protozoans (26.43%) and 33 with *Campylobacter* sp (23.57%) were also observed (Table 1). From the 107 patients with multi-infections, 74 (30.0%) had two or more pathogens; 27 (10.9%) had three or more pathogens; and six

(2.4%) had four pathogens. In patients with two or more pathogens, helminth infections (41.1%) prevailed over protozoan (32.7%) and *Campylobacter* spp. (26.2%) infections.

The most common co-infections were with either *A. lumbricoides* or *Campylobacter* spp. (Table 2). The most prevalent co-infections were *A. lumbricoides* + *E. histolytica*/*E. dispar*, *A. lumbricoides* + *Campylobacter* spp. and *A. lumbricoides* + *T. trichiura*. Total co-infections were highest in 0- to 4- and 5- to 8-yr-old males and were consistently lower throughout the female age groups. The overall prevalence of co-infections in males and females were the same. Of all the co-infections, *A. lumbricoides* was the only one that was significantly associated with the other four pathogens

Table 3. Analysis of pathogens in co-infections of children.

Pathogen 1 vs. pathogen 2	Est. odds ratio	95% CI
<i>Ascaris lumbricoides</i> vs.		
<i>Campylobacter</i> spp.	2.400 ^a	1.282-4.718
<i>Entamoeba histolytica</i> / <i>E. dispar</i>	3.273 ^a	1.630-7.128
<i>Giardia duodenalis</i>	3.286 ^a	1.775-6.472
<i>Trichuris trichiura</i>	10.000 ^a	1.615-38.481
<i>Campylobacter</i> sp. vs.		
<i>Ascaris lumbricoides</i>	0.417 ^a	0.212-0.780
<i>Entamoeba histolytica</i> / <i>E. dispar</i>	1.286	0.700-2.392
<i>Giardia duodenalis</i>	2.083 ^a	1.009-4.551
<i>Trichuris trichiura</i>	2.133 ^a	1.112-4.239
<i>Entamoeba histolytica</i> / <i>E. dispar</i> vs.		
<i>Ascaris lumbricoides</i>	0.306 ^a	0.140-0.614
<i>Campylobacter</i> spp.	0.778	0.418-1.428
<i>Giardia duodenalis</i>	1.438	0.727-2.911
<i>Trichuris trichiura</i>	1.733	0.885-3.520
<i>Giardia duodenalis</i> vs.		
<i>Ascaris lumbricoides</i>	0.304 ^a	0.155-0.563
<i>Campylobacter</i> spp.	0.480 ^a	0.220-0.991
<i>Entamoeba histolytica</i> / <i>E. dispar</i>	0.696	0.343-1.375
<i>Trichuris trichiura</i>	1.286	0.604-2.793
<i>Trichuris trichiura</i> vs.		
<i>Ascaris lumbricoides</i>	0.100 ^a	0.026-0.277
<i>Campylobacter</i> spp.	0.469 ^a	0.236-0.891
<i>Giardia duodenalis</i>	0.778	0.358-1.655
<i>Entamoeba histolytica</i> / <i>E. dispar</i>	0.577	0.284-1.131

Table 4. Prevalence of three pathogens in children.

Table 4: Prevalence of three pathogens in children					
	Number infected (%)				
	Age groups (yrs)				
Male multi-pathogens	0-4	5-8	9-12	Total	
<i>A.l.</i> + <i>T.t.</i> + <i>C.sp.</i>	1 (3.70)	1 (3.70)	0 (0.00)	2 (7.41)	
<i>A.l.</i> + <i>T.t.</i> + <i>E.h.</i>	1 (3.70)	0 (0.00)	0 (0.00)	1 (3.70)	
<i>A.l.</i> + <i>T.t.</i> + <i>G.d.</i>	0 (0.00)	2 (7.41)	0 (0.00)	2 (7.41)	
<i>A.l.</i> + <i>G.d.</i> + <i>C.sp.</i>	1 (3.70)	1 (3.70)	0 (0.00)	2 (7.41)	
<i>A.l.</i> + <i>G.d.</i> + <i>E.h.</i>	0 (0.00)	0 (0.00)	0 (0.00)	0 (0.00)	
<i>A.l.</i> + <i>E.h.</i> + <i>C.sp.</i>	0 (0.00)	1 (3.70)	0 (0.00)	1 (3.70)	
<i>T.t.</i> + <i>G.d.</i> + <i>E.h.</i>	0 (0.00)	1 (3.70)	0 (0.00)	1 (3.70)	
<i>T.t.</i> + <i>G.d.</i> + <i>C.sp.</i>	0 (0.00)	1 (3.70)	0 (0.00)	1 (3.70)	
<i>T.t.</i> + <i>E.h.</i> + <i>C.sp.</i>	0 (0.00)	0 (0.00)	0 (0.00)	0 (0.00)	
<i>G.d.</i> + <i>E.h.</i> + <i>C.sp.</i>	0 (0.00)	0 (0.00)	0 (0.00)	0 (0.00)	
Total	3 (11.11)	7 (25.97)	0 (0.00) ^a	10 (37.04)	
Female multi-pathogens	0-4	5-8	9-12	Total	All
<i>A.l.</i> + <i>T.t.</i> + <i>C.sp.</i>	3 (11.10)	0 (0.00)	1 (3.70)	4 (14.81) ^b	6 (22.22)
<i>A.l.</i> + <i>T.t.</i> + <i>E.h.</i>	0 (0.00)	0 (0.00)	2 (7.41)	2 (7.41)	3 (11.10)
<i>A.l.</i> + <i>T.t.</i> + <i>G.d.</i>	2 (7.41)	0 (0.00)	0 (0.00)	2 (7.41)	4 (14.81)
<i>A.l.</i> + <i>G.d.</i> + <i>C.sp.</i>	1 (3.70)	0 (0.00)	0 (0.00)	1 (3.70)	3 (11.10)
<i>A.l.</i> + <i>G.d.</i> + <i>E.h.</i>	1 (3.70)	0 (0.00)	0 (0.00)	1 (3.70)	1 (3.70)
<i>A.l.</i> + <i>E.h.</i> + <i>C.sp.</i>	0 (0.00)	2 (7.41)	2 (7.41)	4 (14.81) ^b	5 (18.52)
<i>T.t.</i> + <i>G.d.</i> + <i>E.h.</i>	0 (0.00)	0 (0.00)	1 (3.70)	1 (3.70)	2 (7.41)
<i>T.t.</i> + <i>G.d.</i> + <i>C.sp.</i>	0 (0.00)	1 (3.70)	0 (0.00)	1 (3.70)	2 (7.41)
<i>T.t.</i> + <i>E.h.</i> + <i>C.sp.</i>	0 (0.00)	0 (0.00)	0 (0.00)	0 (0.00)	0 (0.00)
<i>G.d.</i> + <i>E.h.</i> + <i>C.sp.</i>	0 (0.00)	1 (3.70)	0 (0.00)	1 (3.70)	1 (3.70)
Total	7 (25.92)	4 (14.81)	6 (22.22)	17 (62.96)	27 (100.00)

(A.l.: *Ascaris lumbricoides*, T.t.: *Trichuris trichiura*, E.h.: *Entamoeba histolytica*/E. *dispar*, G.d.: *Giardia duodenalis*, C.spp.: *Campylobacter* spp.)^aTriple infections were least prevalent in 9-12-yr-old males, P < 0.05.^bA. *lumbricoides* + T. *trichiura* + *Campylobacter* sp. and A. *lumbricoides* + E. *histolytica*/E. *dispar* + *Campylobacter* spp. were significantly higher than other triple infections for females over males, P < 0.05.**Table 5.** Prevalence of four pathogens in children.

		Number infected (%)			
		Age groups (yrs)			
Male multi-pathogens	0-4	5-8	9-12	Total	
<i>A.l.</i> + <i>T.t.</i> + <i>E.h.</i> + <i>C.sp.</i>	0 (0.00)	1 (16.67)	0 (0.00)	0 (0.00)	
<i>A.l.</i> + <i>T.t.</i> + <i>E.h.</i> + <i>G.d.</i>	0 (0.00)	0 (0.00)	0 (0.00)	0 (0.00)	
<i>A.l.</i> + <i>T.t.</i> + <i>G.d.</i> + <i>C.sp.</i>	0 (0.00)	0 (0.00)	0 (0.00)	0 (0.00)	
<i>T.t.</i> + <i>G.d.</i> + <i>E.h.</i> + <i>C.sp.</i>	0 (0.00)	0 (0.00)	0 (0.00)	0 (0.00)	
Total	0 (0.00)	1 (16.67)	0 (0.00)	1(16.67)	
Female multi-pathogens	0-4	5-8	9-12	Total	All
<i>A.l.</i> + <i>T.t.</i> + <i>E.h.</i> + <i>C.sp.</i>	0 (0.00)	1 (16.67)	1 (16.67)	2 (33.33)	3 (50.00)
<i>A.l.</i> + <i>T.t.</i> + <i>E.h.</i> + <i>G.d.</i>	0 (0.00)	0 (0.00)	0 (0.00)	0 (0.00)	0 (0.00)
<i>A.l.</i> + <i>T.t.</i> + <i>G.d.</i> + <i>C.sp.</i>	1 (16.67)	1 (16.67)	0 (0.00)	2 (33.33)	2 (33.33)
<i>T.t.</i> + <i>G.d.</i> + <i>E.h.</i> + <i>C.sp.</i>	0 (0.00)	1 (16.67)	0 (0.00)	1 (16.67)	1 (16.67)
Total	1 (16.67)	3 (50.00) ^a	1 (16.67)	5 (83.33) ^b	6 (100.00)

(A.l.: *Ascaris lumbricoides*, T.t.: *Trichuris trichiura*, E.h.: *Entamoeba histolytica*/E. *dispar*, G.d.: *Giardia duodenalis*, C.spp.: *Campylobacter* spp.)^aFour of the six multi-infections were found in the 5-8-year-old groups.^bFive of the six multi-infections were found in females.

(Table 3). *Campylobacter* spp. was significantly associated with all pathogens except *E. histolytica/E. dispar*. *E. histolytica/E. dispar* + *A. lumbricoides*, *G. duodenalis* + *A. lumbricoides*, *T. trichiura* + *A. lumbricoides* and *T. trichiura* + *Campylobacter* spp. were significantly associated with each other.

Both males and females had more triple infections of *A. lumbricoides* + *T. trichiura* + *Campylobacter* spp. and *A. lumbricoides* + *E. histolytica/E. dispar* + *Campylobacter* spp. than other infection combinations (Table 4). Nine- to twelve-year-old males did not have triple infections. Females had the most infections with four pathogens (Table 5).

Discussion

Children living in both Guatemalan highland and riparian communities are infected soon after weaning and likely re-infected during the rest of their childhood. This prevalence of multi-infections is affected by malnutrition [7], by differences in the behavior of children, by irregular distributions of infecting stages in the environment, by differences in the ability to generate an adequate immunological response, by basic biological differences between parasites, and by host genetic differences [2]. Previous studies have shown that individuals with multiple infections tend to display higher intensities of infection than that expected for each infection separately [8,9]. Thus, infection by one of the pathogens may be influenced by concurrent or earlier infections with the other.

The use of patients with diarrhoeal stools in the absence of controls in the present study limits the analysis of the results of multi-infections associated with patients' exhibiting abdominal symptoms; and the absence of information on the presence of fever, bloody stools, or chronicity of diarrhoea further limits the results to patients with nonspecified abdominal symptoms. However, our study demonstrated that the extent of multi-gastrointestinal pathogen prevalence was different by gender and was strongly associated with age. Infants and young children were more likely to have multiple infections when compared to older children, a finding consistent with lifestyle differences. This coexistence of pathogens in children could have been influenced by the presence of malnutrition or a low immune state resulting in reduced host resistance to infection.

Reported pathogen prevalence rates are not always comparable due to the use of different

research methods, *i.e.*, the use of different diagnostic techniques, or selection of geographical locations, or socioeconomic patterns. Therefore, this high prevalence of co-infections emphasizes the need for more comprehensive analysis of stool samples for pathogens that relate to disease symptoms.

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